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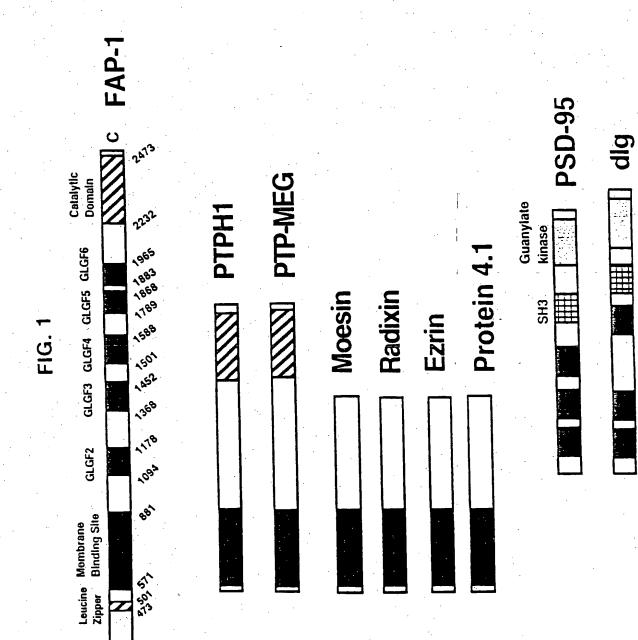
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22

B

GLGF1

Construction of pBTM116 (LexA)-(X)15 DUP16-FAP-1 Library DNAs of pBTM116 (LexA)-(X)15 Large scale transformation of yeast L40 Tehn-(H)15 PUP16-FRP-1 His+, β -gal+ Curing of pVP16-FAP-1 Isolation of pBTM116 (LexA)-(X)15

Analysis of DNA sequences

8-1 9-3

0-5

14-1

DSENSNFRNEIOSLV		SISNSRNENEGGOSLE	STPDTGNENEGGCLE
DSEN	· .	SISN	 STPD
IG. 2B Human		Rat	Mouse

FIG. 2D	
	•
	•
٠	1
•	•

FIG. 2C

IPPDSEDGNEEQSLV	DSEMYNFRSQLASVV	IDLASEFLFLSNSFL	L S	တ	ဟ	<u> </u>	RGFISLV	> %	
12-0	2-0	13-0	20-0	6-5	9-2	18-1	22-1	71-1	14-5
CYA A I G L V	ENA GVS E	WWG ATQ P	E H A O O O	NSS FHS L	GLR LPP D V	GSD SGV N	K K R P V N V	I G K D V W A	ASR NEE L

57-5 72-1 25-9 16-13 6-3

FIG. 3A

Fas C terminal 15 a.a. peptide (μ M)



200 -

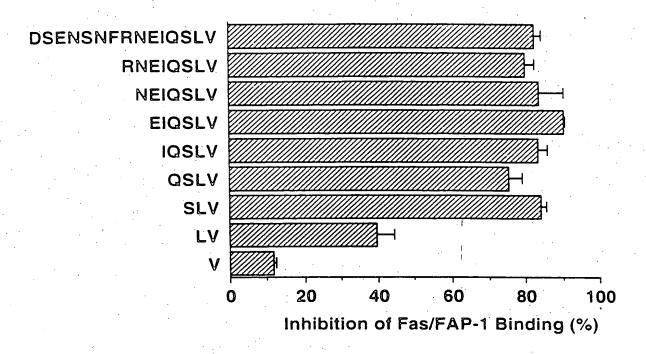
97.4 -FAP-1 69 -

46 -

30 –

21.5 **–** 14.3 **–**

FIG. 3B



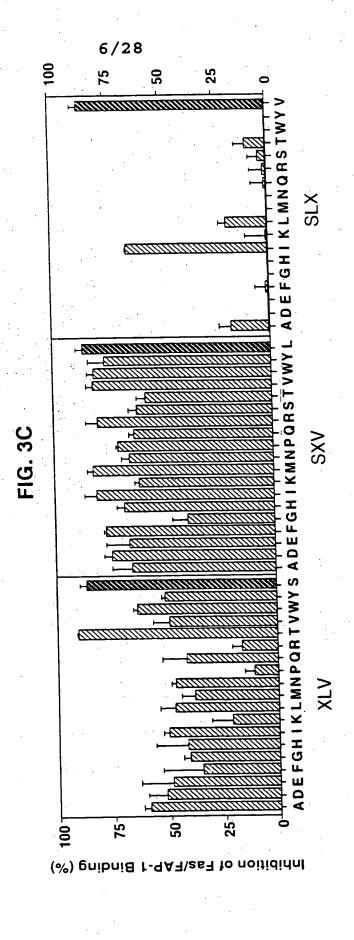


FIG. 4A

VF	716	<u>VP16</u>
FAP-1	Ras	FAP-1 Ras
		多母母 金件等。
0	9000	
	000	TE E GOE
	1000	COSEES
	FAP-1	

His+

His -

FIG. 4B



250 -148 -

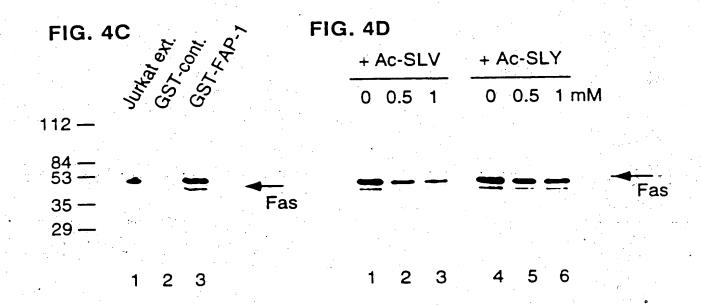
← FAP-1

60 -

42 -

30 -

1 2 3



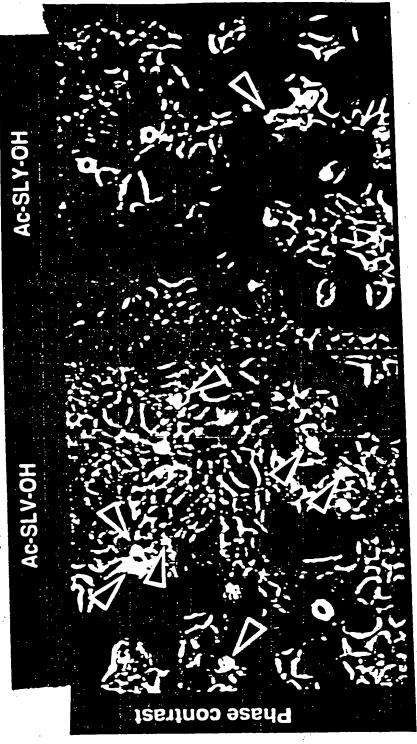
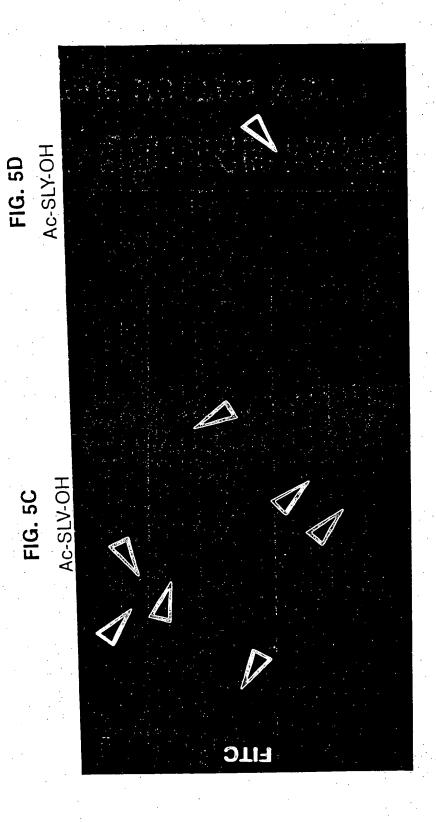


FIG. 5B

FIG. 5A



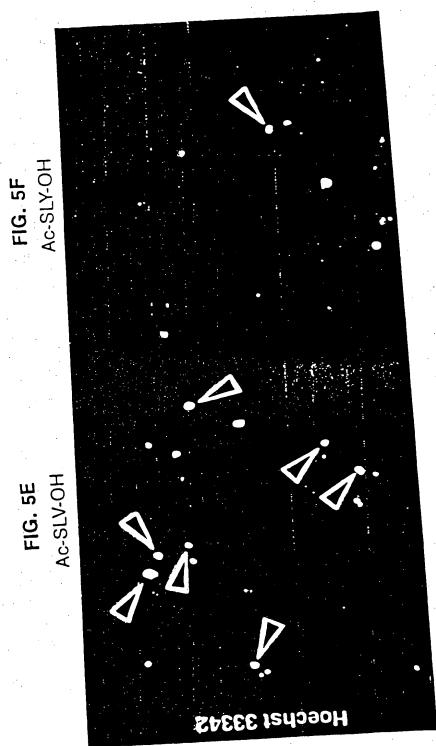


FIG. 5E

FIG. 6

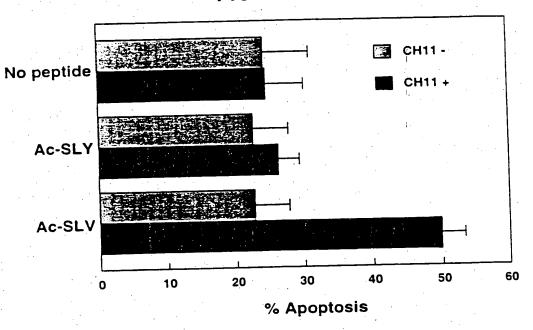


FIG. 7A

Receptor

NGF

edterglrec vvttvmgssq nqtpppegek lngsagdtwr adlveslcse ygyyddettg egvaqpcgan eaddavcrca qdliastvag nkqgansrpv pakreevekl gecckacnlg vdpclpctvc llaalrrigr acptglyths gldsmsapcv dgtysdeanh ddsslysslp afkrwnsckg stdepeappe atqdsatlda lgvslggake epckpctecv cpvrallasw kgntvceecp ppegsdstap tqtasgqalk avvvglvayi lipvycsila sqslhdqqph ehidsfthea gsglvfscqd ipgrwitrst dgprllllll vtfsdvvsat pvvtrgttdn mgagatgram qtvcepclds **lhsdsgisvd** hlagelgyqp rceacrycea trwadaecee stat**spv** 421 361 181 241 301

FIG. 7B

CD4 Receptor

yagsgnltla kltgsgelww vskrekavwv vsqlelqdsg figlgiffcv fhwknsngik vedgkeevgl iedsdtyice sfplaftve hltlpgalpg ctasqkksiq kniqggktls slklenkeak vlggvaglll llvlqlallp aatqgkkvvl gkkgdtvelt nfpliiknlk vykkegegve stpvqpmali psydcrsprg klqmgkklpl qlqknltcev wgptspklml faktcspi ekktcgcphr dsrrslwdqg vsvkrvtqdp esnikvlptw ltlesppgss vlafqkassi rmsgikrlls kgpsklndra kkvefkidiv witfdlknke thllqgqslt evnlvvmrat 11sdsgqv11rcrhrrrgae twtctvlqnq gaerasssks Inpeagmwdc ilgnqgsflt lvfgltansd leaktgklhq mnrgvpfrhl 361

FIG. 7C

Species	C-terminal sequences of NGFR (p75)	Binding activity of FAP-1
Human	SESTATSPV-COOH	+
Rat	SESTATSPV-COOH	+
Chicken	SESTATSPV-COOH	+

FIG. 70

rttcsenela nlvaayekak agcsvqpwes wekelagire gpsspgrits les!gvsssv sibidplsy streageday kklakaqceq selrselsgs nvvcgrkkss segnalltit slilgqfraa dgecggafav fundlkrans qtererdlle ssdrpvlgse llalaesed elgrvitgle qrrlqsvqat kekkalelkl ypnlaeersr eringrichi gttireedey aavkitmlel aspalelael rialleeens seirhqqsae ysagcieaye dyiqq1)mdr sikaqiyile dkpgkecada divelnkrlq elnkkidrlg dadacsdins clsktreess skirefevet elegvlgrdl naakallmkl drlrrrvrel hetqvrmlkg vsaleritks slsstssgsk lyshgsalse rahdcrktae elmamkeema plakiaerv natalrialq ftkedegr1k hcdlaiktve ritelhsvia pengetmyta hsaalaslkg armaireer esquamver gdenitamlk msmlvgkyes sstasscdte qerttlryee hieglttase csniqeifqt ldlenavlmq veedkagrmr kklkarvqel elstssssnd mddddtsvs1 skaaalnrtk ndssaelsel aeftnairre lvhiehlkse ach ahslqd rygsepgdas senahtstt avkprgdsgr enesitamic nrpinpstg shlmrehedv ghevnedsra eecksnaer msgvamkyg cslsvaevdr 661 301 361 421 601 181 241 481 541

FIG. 7E

pegdeegrame gkvmladrkg mlagqppfdg spefedhegs drlyfvmeyv ldseghikia vrehaffrri sdfeafsyvn shctdfiwgf eklhvtvrda kpsdkdrrl waygvllye lgcgpegerd dalvianidq arffkoptfc riylkaevad pownesftfk 1mvlgkgsf glhscfqtv yrdlkldnvm khkfklhtyg egeyynvpi ergapy1tpp kgpdtddprs ayqpygkavd glmtkbpakr candhtekrg ktktirstin vhevkdhkf1 sgwykllng drvkltdfn allddopfl. Lfflhkrgii vtfscpgad papameska. Egvselmkmp vectimekry vfyaaeisig pdyiapeii lskeavs1ck kgaenfdkff rkgalrdm sedrkgpsnn kacvinvpsl mehnysypks tradfmgsls gstvytage ckdvvigddd qvgkfkepqa dgvttrtfcg appfkpkvcg cfvvhkrche kedtedmnyh **casqdvanr**f sdpyvk1k1 dweklenre1 adadelfqsi madvfpgnds gkagfacave nggdlmybiq [fgmckehum lyglihqgm inlipmodphg sveiwdwdrt eelyaikil rokfekaki

FIG. 7F

[dryva; qnp seklfgrsih escnedvíga vntipalayk yflmslaiad ddnfvligef nrtalscego sdgvnekvag v enkkplqlil itrimavick stipqsele asimpleais vfkegsclla dafnwtvdse lekklqnatn qhseeaskdn fsrylqcqyk gnilvimavs pvfglgddsk digtraklas 1 formwepff dfnsgearts wiyldvlfst tisvgismpi iqkeaticvs ndcsmvalgk tlfnktyrsa wplpsklcav tavviiltia ackvlgivff 1nddtrlysn lestinslmd lgeknwsall lssavnplvy vityfltiks nskadaktid tmgs1sneqk kaflkilavw smitilygyr lspsclallh mllgflympv vsffipleim 11nvfvwigy seglangakk epgaytgrr md:1ceents hinsrfngrt

FIG. 7G

gdkteeqwex dscnqttlgm reakiyfrnp cafikituw asmivtyflt kalpnøgdet alltinfeam gnklhwaall dtllltene atkevktlrk eemkqiveed llvglfvmpf lgangynera vamldgsrkd fithitivic slaafftpl wagldtesip yflmelavad vdryiaikkp kerfgdfmlž detpcsspak Elflimopf foryitemyr raskvlgivf tl£nktfrda npnniccv1t lekklqyatn asimhleais witvetyfor stfvhvissn gmtlvilavs wlfldvlfst pikgietdvd vknkppqrlt kevatiened vasgvnplvy hgirnginpa asigiaipv maenskffkk wplplvlcpa mrrtstigk Lleifwigy ilmvilptig halqkkayl malsyrvsel 121 181 241 301 361 421

FIG. 7H

```
i maaasydgii kqvealkmen snirqeledn snhitklete asnmkevikq iqqsiedeam
 1 assgçidile rikelnidss nfpgvklrsk msirsygere gsvssrsged spvpmgsfpr
121 rgfvngsres tgyleeleka rsllladldk eekekdwyya qlqmltkrid slpltenfsi
181 qtdmtrrqle yearqirvam eeqlqtcqdm ekraqrriar iqqiekdilr irqllqsqat
241 eaerssqukh eigshdaerg negggvçein matagngggs tirmdhetas vissssthsa
301 przitahlgt kvemvyslis migthdkódm sztliamses gdscismrgs golpiliqli
361 hgndkdsvil gnsrgskear arasaalhni ihsopddkrg rzeirvihli eqiraycsto
421 wewgeahepg mdqddcnpmpa pvehgicpay cylmkiside emmamnelg glqaiaellq
461 vdcemygltm dhysitlrry agmaltnitf gdvamkatlo smkgcmralv aqlksesedi
541 qqviasvlrn lswradvnsk ktlrevgsvk almecalevk kestlksvls alwnisahc:
601 enkadicavd galaflygtl tyrsqtntla iiesgggilr nyssliatne dhrqilrenn
661 clqtllqhlk shsltivsna cqtlwnlsar npkdqaalwd mgavsmlknl ihskhkmiam
721 gsaaairnim anrpakykda nimspyssip sihvrkokal eaeldaghis etfonionis
781 pkashrskqr hkqslygdyv fdtnrhddnr sdnfntgnmt vispylnttv ipssaasrqs
 841 ldasraekdr slerergigl gnyhpateno gtaskralqi sttaaqiakv meevsaihta
901 qedrssgatt elhovtdern alrrasaaht hantynitka eranrtosmo yakleykras
961 ndslnavses dgygkrgqmk psiesysedd eskfcsygqy padlabkihs arbmddndge
1021 ldcpinyslk ysdeglnagr gspagnerwa rpkhiledei kqaeqrqarn qsttypvyte
1081 stddkhlkfq phfgqqecvs pyrsrgangs etnrvgsnig inqnvsqslc qeddyeddkp
1141 tnyserysee eqheeeerpt nysikyneek rhvdqpidys lkyatdipss qkqsfsfsks
1201 saggsakteh messsentst pasnakrong lhpssagsrs gopokaatek vasingetig
1261 tycvedtpic fsrcsslasi ssaedeigen qttqeadsan tiqiaeikek igtrsaedpv
1321 sevpaysonp rikssriggs sissesarhk avefssgaks paksgagtpk sppehyvget
1381 plmfarctsv ssldsfesrs iassvqsepc sgmvsgilsp sdlpdspgqt mppsraktpp
1441 pppqtaqtkr evpknkapta ekresgpkqa avnaavqrvq vlpdadtilh fatestpdgf
1501 scassisals idepfickdy elrimppyge ndngmetese opkesnenge keaektidse
1561 kdilddøddd dieileecii samptkeerk akkpaqtask lpppvarkps glpvyklips
1621 qurlqpqkhv sftpqddmpr vycvegtpin fstatsledi tiesppnela agegvrggaq
1681 sgefekrdti pregrstdea gggktssvii pelddnkaee gdilaecins ampkgkshkp
1741 frykkindav agasessap næglagkkk ketspykeie anteyrtryr knadskanla
1801 aervisdikd skkonlknns kdindklpnn edryrgsiai dsprhytpie gtpycisrnd
1861 slasldfddd dydlsrekae irkakenkes eakytshtel tanggankt galakopinr
 1921 gapkpilaka stipasski pargastdek lanfalentp voishnssls sladidaenn
 1981 nkenepiket eppdaggeps kpqasgyapk sfhvedtpvc fsrasslssi sidseddllq
 2041 ecissampkk kkpsrikgdn ekhspringg ilgeditldl kdigipdseh glspdsenfd
 2101 wkaigegans ivssihqaaa aacisrqass dsdsiislks gislgspfhl tpdqeexpft
 2161 ankaprilkp gekstletkk ieseskgikg gkkvykslit gkvrsnseis ggmkaplaan
 2221 mpsisrgrtm ihipgvrnss sstspyskkg pplktpasks pseggtatts prgakpsyks
 2281 elapvarque qiggeskape regerdetpe rpacquelerp iceporneis pgrngispon
 2341 klsqlprtss pstastkssg sgkmaytspg romsqqzltk qtglskmass iprsesaskg
 2401 lnqmnngnga nkkvelsrms stkssgsesd rserpvlvrq stfikeapsp tlrrkleesa
 2461 efeslapesr pasptragad tpvlspslpd malathaevd aggwrklppm leptieyndg
 2521 rpakridiar shaesparly inragtwkre hakhasalpr vatwrrtgas asilsasses
2531 sekaksedek hynaisgtko skenovako twrkikenef aptnatagtv asgatngaes
 2541 ktliyqmapa vsktedvwvr iedopinnpr sgrsptgntp pvidsvseka npnikdskdn
 2701 qakqnvgngs vpmrtvglen rlnsfiqvda pdqkgteikp gqnnpvpvse tnessivert
 2761 pissesskh espegivaar vipinynpep rkssadsisa rpsqipipvm nnikkrdekt
 2621 datessgtqs pkrhagaylv ter
```

FIG. 8

(Low-affinity nerve growth factor receptor) p75NGFR

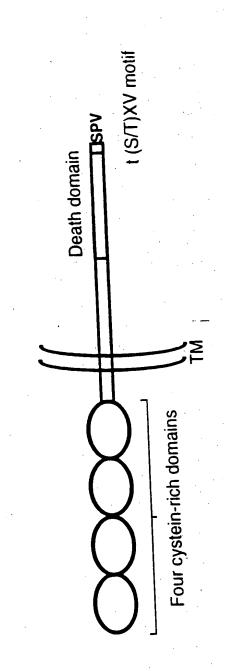


FIG. 9

C-terminal amino acid sequence	NEIQSLV	STATSPV
	Fas	p75NGFR

◆ → PDZ domain

t (S/T)-X-V

interaction

FIG. 10

In vitro interaction of 35S-labeled FAP-1 with various receptors FAP-1 binds to the cytoplasmic region of p75NGFR.

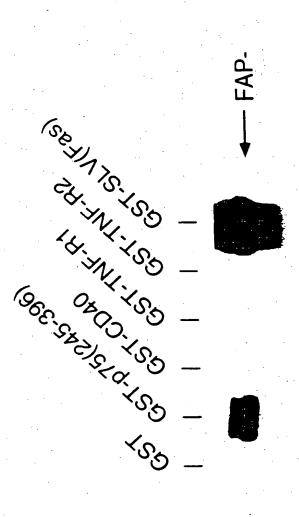
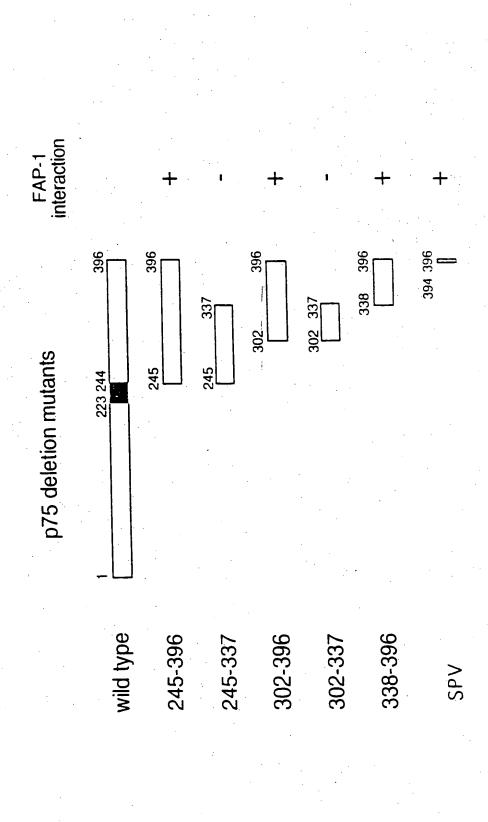


FIG. 11A

FAP-1 binds to C-terminal three amino acids SPV of p75NGFR.



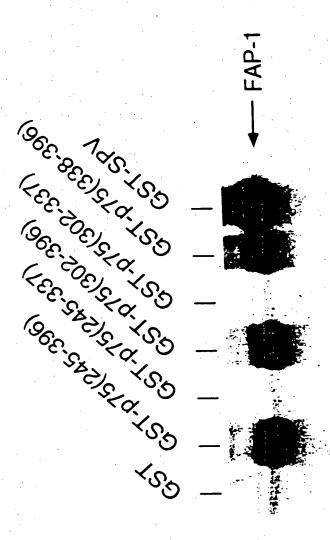


FIG. 11B

FIG. 12

FAP-1 binds to p75NGFR C-terminal cytoplasmic region in yeast

	VP16-FAP-1	VP16-cRaf
_exA-p75NGFR(338-396)	+	•
LexA-p75NGFR(365-396)	+	1
LexA-Fas	+++	1
LexA-Ras ^{V12}		+
LexA-Lamin	•	•

FIG. 13A

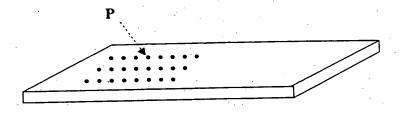


FIG. 13B

(S/T)-X-(V/I/L)-COOH

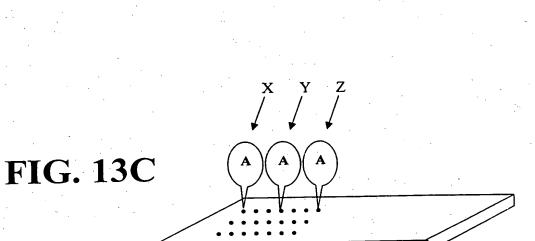


FIG. 14A Plain-glass slide FIG. 14B 3D gel pad chip Selection of the s